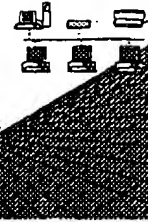


CRF

BIOTECHNOLOGY  
SYSTEMS  
BRANCH

SEP 11 2003



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/786,502  
Source: 1600  
Date Processed by STIC: 9-9-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)

2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450

3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name,  
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

Or

U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,  
2011 South Clark Place, Arlington, VA 22202

4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,  
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

# Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/786,502
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 ____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 ____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 ____ Misaligned Amino Numbering	The numbering under each 5 <sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 ____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 ____ Variable Length	Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 ____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 ____ Skipped Sequences (OLD RULES)	Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped  Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 ____ Skipped Sequences (NEW RULES)	Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 ____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 ✓ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 ____ Use of <220>	Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 ____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 ____ Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.	

#16  
RTH 7/24/03Does Not Comply  
Corrected Diskette Needed

1600

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003

TIME: 07:00:07

Input Set : N:\efs\09786502\ptoms.txt

Output Set: N:\CRF4\09092003\I786502.raw

3 <110> APPLICANT: Sadelain, Michel  
 4 Bander, Neil H  
 5 Gong, Michael  
 7 <120> TITLE OF INVENTION: Fusion Receptors Specific For Prostate-Specific Membrane  
 Antigen and Uses  
 8 Thereof  
 10 <130> FILE REFERENCE: MSK.P-040  
 12 <140> CURRENT APPLICATION NUMBER: 09/786,502  
 13 <141> CURRENT FILING DATE: 2001-05-18  
 15 <150> PRIOR APPLICATION NUMBER: PCT/US99/20349  
 16 <151> PRIOR FILING DATE: 1999-09-03  
 18 <150> PRIOR APPLICATION NUMBER: US 60/099,138  
 19 <151> PRIOR FILING DATE: 1998-09-04  
 21 <160> NUMBER OF SEQ ID NOS: 8  
 23 <170> SOFTWARE: PatentIn version 3.1  
 25 <210> SEQ ID NO: 1  
 26 <211> LENGTH: 22  
 27 <212> TYPE: DNA  
 28 <213> ORGANISM: synthetic construct  
 30 <400> SEQUENCE: 1  
 31 aggtsmarct gcagsagtcw gg 22  
 34 <210> SEQ ID NO: 2  
 35 <211> LENGTH: 34  
 36 <212> TYPE: DNA  
 37 <213> ORGANISM: synthetic construct  
 39 <400> SEQUENCE: 2  
 40 tgaggagacg gtgaccgtgg tcccttgccc ccag 34  
 43 <210> SEQ ID NO: 3  
 44 <211> LENGTH: 24  
 45 <212> TYPE: DNA  
 46 <213> ORGANISM: synthetic construct  
 48 <400> SEQUENCE: 3  
 49 gacattgagc tcaccagtc tcca 24  
 52 <210> SEQ ID NO: 4  
 53 <211> LENGTH: 33  
 54 <212> TYPE: DNA  
 55 <213> ORGANISM: synthetic construct  
 57 <400> SEQUENCE: 4  
 58 tgcggccgcc cgtttgatct ccagcttggt ccc 33  
 61 <210> SEQ ID NO: 5  
 62 <211> LENGTH: 43  
 63 <212> TYPE: DNA  
 64 <213> ORGANISM: synthetic construct  
 66 <400> SEQUENCE: 5

See item #10 on ERROR  
Summary SHEET.

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003

TIME: 07:00:07

Input Set : N:\efs\09786502\ptoms.txt

Output Set: N:\CRF4\09092003\I786502.raw

67 gaagaagatc tgacattgtg atgaccagtc tcacaaattc atg 43  
70 <210> SEQ ID NO: 6  
71 <211> LENGTH: 40  
72 <212> TYPE: DNA  
73 <213> ORGANISM: synthetic construct  
75 <400> SEQUENCE: 6  
76 tgcggccgcc cgtttcaggt ccagcatggt cccagcaccg 40  
79 <210> SEQ ID NO: 7  
80 <211> LENGTH: 27  
81 <212> TYPE: DNA  
82 <213> ORGANISM: synthetic construct  
84 <400> SEQUENCE: 7  
85 gcggccgcaa ttgaagttat gtatcct 27  
88 <210> SEQ ID NO: 8  
89 <211> LENGTH: 30  
90 <212> TYPE: DNA  
91 <213> ORGANISM: synthetic construct  
93 <400> SEQUENCE: 8  
94 tcgaggatct tgtcaggagc gataggctgc 30

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003  
TIME: 07:00:08

Input Set : N:\efs\09786502\ptoms.txt  
Output Set: N:\CRF4\09092003\I786502.raw

Invalid Line Length: *OK*

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/786,502

DATE: 09/09/2003

TIME: 07:00:08

Input Set : N:\efs\09786502\ptoms.txt

Output Set: N:\CRF4\09092003\I786502.raw